

**RAW SEQUENCE LISTING
ERROR REPORT**

BEST AVAILABLE COPY

BIOTECHNOLOGY
SYSTEMS
BRANCH

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/700,643

Source: PC909

Date Processed by STIC: 1/11/2001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin30help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO).

Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

PCT09

as C1107 once

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/700,643

DATE: 01/13/2003
TIME: 17:18:71

Input Set : A:\2523.sequence.list.txt
Output Set: H:\CRF3\01112001\I700643.raw

5 <110> APPLICANT: Matsumoto, Hirokazu
W--> 7 <110> APPLICANT: Kitada, Chieko
W--> 9 <110> APPLICANT: Hinuma, Shuji
11 <120> TITLE OF INVENTION: ANTI BODY AND USE THEREOF
13 <130> FILE REFERENCE: 25130506
C--> 15 <140> CURRENT APPLICATION NUMBER: US/09/700,643
C--> 15 <141> CURRENT FILING DATE: 2000-11-16
15 <150> PRIORITY APPLICATION NUMBER: PCT/JP99/02650
W--> 17 <150> PRIORITY APPLICATION NUMBER: 1999-05-20
W--> 19 <150> PRIORITY APPLICATION NUMBER: JP 10-140293
21 <151> PRIORITY FILING DATE: 1998-5-21
23 <160> NUMBER OF SEQ ID NOS: 13
27 <210> SEQ ID NO: 1
29 <211> LENGTH: 31
31 <212> TYPE: PRT
33 <213> ORGANISM: Bovine
35 <223> OTHER INFORMATION: Xaa means Phe-NH2
37 <400> SEQUENCE: 1
39 Ser Arg Ala His Glu His Ser Met Glu Thr Arg Thr Pro Asp Thr Asn
40 5 10 15
OK> 42 Pro Ala Trp Tyr Ala Gly Arg Gly Ile Arg Pro Val Gly Arg Xaa
43 20 25 30
46 <210> SEQ ID NO: 2
48 <211> LENGTH: 31
50 <212> TYPE: PRT
52 <213> ORGANISM: Human
54 <223> OTHER INFORMATION: Xaa means Phe-NH2
56 <400> SEQUENCE: 2
58 Ser Arg Thr His Arg His Ser Met Glu Ile Arg Thr Pro Asp Ile Asn
59 5 10 15
OK> 61 Pro Ala Trp Tyr Ala Ser Arg Gly Ile Arg Pro Val Gly Arg Xaa
62 20 25 30
65 <210> SEQ ID NO: 3
67 <211> LENGTH: 31
69 <212> TYPE: PRT
71 <213> ORGANISM: Rat
73 <223> OTHER INFORMATION: Xaa means Phe-NH2
75 <400> SEQUENCE: 3
77 Ser Arg Ala His Glu His Ser Met Glu Thr Arg Thr Pro Asp Ile Asn
78 5 10 15
OK> 80 Pro Ala Trp Tyr Thr Gly Arg Gly Ile Arg Pro Val Gly Arg Xaa
81 20 25 30
84 <210> SEQ ID NO: 4
86 <211> LENGTH: 15
88 <212> TYPE: PRT
90 <213> ORGANISM: Human
92 <223> OTHER INFORMATION: Xaa means Phe-NH2

Does Not Comply
Corrected Diskette Needed

ppr 1-3

use C1517 for PRIORITY FILING DATE

Per 1.823 of new sequence rules,

C12207 <insert this mandatory
numerical identifier
whenever C2217,

C2227, or C2237

<insert C2207
(global error)

insert C2207

insert C2207

insert C2207

OK

OK

OK

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/700,643

DATE: 01/11/2001
TIME: 17:18:11

Input Set : A:\2523.sequence.list.txt
Output Set: N:\CRF3\01112001\I700643.raw

OK > 94 <400> SEQUENCE: 4
 96 Pro Ala Trp Tyr Ala Ser Arg Gly Ile Arg Pro Val Gly Arg Xaa
 97 5 10 15
 100 <210> SEQ ID NO: 5
 102 <211> LENGTH: 20
 104 <212> TYPE: PRT
 106 <213> ORGANISM: Human
 108 <223> OTHER INFORMATION: Xaa means Phe-NH2
 110 <400> SEQUENCE: 5
 112 Thr Pro Asp Ile Asn Pro Ala Trp Tyr Ala Ser Arg Gly Ile Arg Pro
 113 5 10 15
 OK > 115 Val Gly Arg Xaa
 116 20
 119 <210> SEQ ID NO: 6
 121 <211> LENGTH: 15
 123 <212> TYPE: PRT
 125 <213> ORGANISM: Human
 127 <223> OTHER INFORMATION: Xaa on the 1st position means Ac-Pro, Xaa on the 4th position means Tyr(AC),
 and Xaa on the 15th position means Phe-NH2
 OK > 129 <400> SEQUENCE: 6
 131 Xaa Ala Trp Xaa Ala Ser Arg Gly Ile Arg Pro Val Gly Arg Xaa
 132 5 10 15
 135 <210> SEQ ID NO: 7
 137 <211> LENGTH: 15
 139 <212> TYPE: PRT
 141 <213> ORGANISM: Human
 143 <223> OTHER INFORMATION: Xaa means Phe-NH2
 145 <400> SEQUENCE: 7
 OK > 147 Cys Ala Trp Tyr Ala Ser Arg Gly Ile Arg Pro Val Gly Arg Xaa
 148 5 10 15
 151 <210> SEQ ID NO: 8
 153 <211> LENGTH: 33
 155 <212> TYPE: PRT
 157 <213> ORGANISM: Bovine
 159 <400> SEQUENCE: 8
 161 Ser Arg Ala His Gin His Ser Met Glu Ile Arg Thr Pro Asp Ile Asn
 162 5 10 15
 164 Pro Ala Trp Tyr Ala Gly Arg Gly Ile Arg Pro Val Gly Arg Phe Gly
 165 20 25 30
 167 Arg
 170 <210> SEQ ID NO: 9
 172 <211> LENGTH: 31
 174 <212> TYPE: PRT
 176 <213> ORGANISM: Human
 178 <223> OTHER INFORMATION: Xaa means Phe-NH2
 180 <400> SEQUENCE: 9
 182 Ser Arg Ala His Gin His Ser Met Glu Ile Arg Thr Pro Asp Ile Asn
 183 5 10 15
 185 Pro Ala Trp Tyr Ala Gly Arg Gly Ile Arg Pro Val Gly Arg Phe
 186 20 25 30

insert 12207

12207

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/700,643

DATE: 01/11/2001
TIME: 17:18:11

Input Set : A:\2523.sequence.list.txt
Output Set: N:\CRF3\01112001\1700643.raw

189 <210> SEQ ID NO: 10
191 <211> LENGTH: 30
193 <212> TYPE: PRT
195 <213> ORGANISM: Human
197 <223> OTHER INFORMATION: Xaa means Arg-NH2
199 <400> SEQUENCE: 10
201 Ser Arg Thr His Arg His Ser Met Glu Ile Arg Thr Pro Asp Ile Asn
202 5 10 15
OK> 204 Pro Ala Trp Tyr Ala Ser Arg Gly Ile Arg Pro Val Gly Xaa
205 20 25 30

208 <210> SEQ ID NO: 11
210 <211> LENGTH: 14
212 <212> TYPE: PRT
214 <213> ORGANISM: Human
216 <223> OTHER INFORMATION: Xaa means Cys-NH2
218 <400> SEQUENCE: 11

OK> 220 Thr Pro Asp Ile Asn Pro Ala Trp Tyr Ala Ser Arg Gly Xaa
221 5 10

224 <210> SEQ ID NO: 12
226 <211> LENGTH: 20
228 <212> TYPE: PRT
230 <213> ORGANISM: Bovine
232 <223> OTHER INFORMATION: Xaa means Phe-NH2
234 <400> SEQUENCE: 12
236 Thr Pro Asp Ile Asn Pro Ala Trp Tyr Ala Gly Arg Gly Ile Arg Pro
237 5 10 15

OK> 239 Val Gly Arg Xaa
240 20

243 <210> SEQ ID NO: 13
245 <211> LENGTH: 20
247 <212> TYPE: PRT
249 <213> ORGANISM: Rat
251 <223> OTHER INFORMATION: Xaa means Phe-NH2
253 <400> SEQUENCE: 13
255 Thr Pro Asp Ile Asn Pro Ala Trp Tyr Thr Gly Arg Gly Ile Arg Pro
256 5 10 15

OK> 258 Val Gly Arg Xaa
259 20

VERIFICATION SUMMARY
 PATENT APPLICATION: US/09/700,643

DATE: 01/11/2001
 TIME: 17:18:12

Input Set : A:\2523.sequenceList.txt
 Output Set: N:\CRF3\01112001\I700643.raw

L:7 M:280 W: Numeric Identifier already exists. <110> found multiple times
 L:9 M:280 W: Numeric identifier already exists. <110> found multiple times
 L:15 M:270 C: Current Application Number differs, Replaced Current Application No
 L:15 M:271 C: Current Filing date differs, Replaced Current Filing Date
 L:17 M:289 W: Numeric Identifier Missing or Out-Of-Order, <150> PRIOR APP FILING DATE
 L:19 M:289 W: Numeric Identifier Missing or Out-Of-Order, <150> PRIOR APP FILING DATE
 L:21 M:356 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
 L:42 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
 L:43 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:1
 L:44 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:1
 L:45 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:1
 L:61 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2
 L:61 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:2
 L:61 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:2
 L:61 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:2
 L:80 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
 L:80 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:3
 L:80 M:259 W: Mandatory Feature missing, <222> not found for SEQ ID#:3
 L:80 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:3
 L:96 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
 L:96 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:4
 L:96 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:4
 L:96 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:4
 L:115 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:5
 L:115 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:5
 L:115 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:5
 L:115 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:5
 L:131 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:6
 L:131 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:6
 L:131 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:6
 L:131 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:6
 L:147 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:7
 L:147 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:7
 L:147 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:7
 L:147 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:7
 L:204 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:10
 L:204 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:10
 L:204 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:10
 L:204 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:10
 L:220 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:11
 L:220 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:11
 L:220 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:11
 L:220 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:11
 L:239 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:12
 L:239 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:12
 L:239 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:12
 L:239 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:12
 L:258 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:13

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/700,643

DATE: 01/11/2001
TIME: 17:18:12

Input Set : A:\2523.sequenceclist.txt
Output Set: N:\CRF3\01112001\1700643.raw

L:258 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#13
L:258 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#13
L:258 M:340 W: (40) "n" or "Xaa" used: Feature required, for SEQ ID#13